

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/551,504
Source: PLT
Date Processed by STIC: 10/14/2005

ENTERED



PCT

RAW SEQUENCE LISTING

DATE: 10/14/2005

PATENT APPLICATION: US/10/551,504

TIME: 10:55:55

Input Set : A:\14875-153US1sq.txt

Output Set : N:\CRF4\10142005\J551504.raw

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3 <110> APPLICANT: Tsunoda, Hiroyuki
4     Nakano, Kiyotaka
5     Orita, Tetsuro
6     Tsuchiya, Masayuki
7     Hirata, Yuichi
9 <120> TITLE OF INVENTION: ANTI-MPL ANTIBODIES
11 <130> FILE REFERENCE: 14875-153US1
C--> 13 <140> CURRENT APPLICATION NUMBER: US/10/551,504
C--> 13 <141> CURRENT FILING DATE: 2005-09-29
13 <150> PRIOR APPLICATION NUMBER: PCT/JP2004/018506
14 <151> PRIOR FILING DATE: 2004-10-12
16 <150> PRIOR APPLICATION NUMBER: JP 2003-415746
17 <151> PRIOR FILING DATE: 2003-12-12
19 <150> PRIOR APPLICATION NUMBER: JP 2004-71763
20 <151> PRIOR FILING DATE: 2004-03-12
22 <150> PRIOR APPLICATION NUMBER: JP 2004-248323
23 <151> PRIOR FILING DATE: 2004-08-27
25 <160> NUMBER OF SEQ ID NOS: 308
27 <170> SOFTWARE: PatentIn version 3.1
29 <210> SEQ ID NO: 1
30 <211> LENGTH: 1572
31 <212> TYPE: DNA
32 <213> ORGANISM: Homo sapiens
34 <400> SEQUENCE: 1
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37 gtgcagctgg tgcagtctgg acctgagggtg aagaagcctg gggcctcagt gaaggtctcc      120
39 tgcaaggctt ctggatacac cttcaccaac tcctggatga actgggtgag gcagaggcct      180
41 ggaaagggtc ttgagtggat gggacggatt tatcctggag atggagaaac tatctacaat      240
43 gggaaattca gggtcagagt cacgattacc gcggaacgat ccacgagcac agcctacatg      300
45 gagctgagca gcctgagatc tgaggacacg gccgtgtatt actgtgagag aggctatgat      360
47 gattactcgt ttgcttactg gggccaggga accacgggtca ccgtctcttc aggtggtggt      420
49 ggatccggag gtggtggatc ggggtggtgga ggatccgata ttgtgatgac tcagctgca      480
51 ctctccctgc ccgtcacccc tggagagccg gcctccatct cctgcaggtc tagtaagagt      540
53 ctctctgata gtaatggcaa cacttacttg tattggttcc agcagaagcc agggcagctc      600
55 ccacagctcc tgatctatcg gatgtccaac cttgcctcag gggtcctga caggttcagt      660
57 ggcagtgatg caggcacagc ttttactactg aaaatcagca gagtggaggc tgaggatggt      720
59 ggggtttatt actgcatgca acatatagaa tatectttta cgttcggcca agggacaaa      780
61 ctggaaatca aaggaggtgg tggatcgggt ggtggtggtt cgggagggcg tggatcgcag      840
63 gtgcagctgg tgcagtctgg acctgagggtg aagaagcctg gggcctcagt gaaggtctcc      900
65 tgcaaggctt ctggatacac cttcaccaac tcctggatga actgggtgag gcagaggcct      960
67 ggaaagggtc ttgagtggat gggacggatt tatcctggag atggagaaac tatctacaat     1020
69 gggaaattca gggtcagagt cacgattacc gcggaacgat ccacgagcac agcctacatg     1080
71 gagctgagca gcctgagatc tgaggacacg gccgtgtatt actgtgagag aggctatgat     1140

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73 gattactcgt ttgcttactg gggccagggg accacgggtca ccgtctcttc aggtgggtgt 1200
75 ggatccggag gtggtggatc ggggtgggga ggatcggata ttgtgatgac tcagtctgca 1260
77 ctctccctgc ccgtcacccc tggagagccg gcctccatct cctgcaggtc tagtaagagt 1320
79 ctctctgata gtaatggcaa cacttacttg tattggttcc agcagaagcc agggcagtct 1380
81 ccacagctcc tgatctatcg gatgtccaac cttgcctcag gggtcctga caggttcagt 1440
83 ggcagtggat caggcacagc ttttactctg aaaatcagca gagtggaggc tgaggatgtt 1500
85 ggggtttatt actgcatgca acatatagaa tatcctttta cgttcggcca agggacccaaa 1560
87 ctggaaatca aa 1572
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91 <211> LENGTH: 524
92 <212> TYPE: PRT
93 <213> ORGANISM: Homo sapiens
95 <400> SEQUENCE: 2
97 Met Asp Trp Thr Trp Arg Phe Leu Phe Val Val Ala Ala Ala Thr Gly
98 1 5 10 15
101 Val Gln Ser Gln Val Gln Leu Val Gln Ser Gly Pro Glu Val Lys Lys
102 20 25 30
105 Pro Gly Ala Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe
106 35 40 45
109 Thr Asn Ser Trp Met Asn Trp Val Arg Gln Arg Pro Gly Lys Gly Leu
110 50 55 60
113 Glu Trp Met Gly Arg Ile Tyr Pro Gly Asp Gly Glu Thr Ile Tyr Asn
114 65 70 75 80
117 Gly Lys Phe Arg Val Arg Val Thr Ile Thr Ala Asp Glu Ser Thr Ser
118 85 90 95
121 Thr Ala Tyr Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val
122 100 105 110
125 Tyr Tyr Cys Ala Arg Gly Tyr Asp Asp Tyr Ser Phe Ala Tyr Trp Gly
126 115 120 125
129 Gln Gly Thr Thr Val Thr Val Ser Ser Gly Gly Gly Gly Ser Gly Gly
130 130 135 140
133 Gly Gly Ser Gly Gly Gly Gly Ser Asp Ile Val Met Thr Gln Ser Ala
134 145 150 155 160
137 Leu Ser Leu Pro Val Thr Pro Gly Glu Pro Ala Ser Ile Ser Cys Arg
138 165 170 175
141 Ser Ser Lys Ser Leu Leu His Ser Asn Gly Asn Thr Tyr Leu Tyr Trp
142 180 185 190
145 Phe Gln Gln Lys Pro Gly Gln Ser Pro Gln Leu Leu Ile Tyr Arg Met
146 195 200 205
149 Ser Asn Leu Ala Ser Gly Val Pro Asp Arg Phe Ser Gly Ser Gly Ser
150 210 215 220
153 Gly Thr Ala Phe Thr Leu Lys Ile Ser Arg Val Glu Ala Glu Asp Val
154 225 230 235 240
157 Gly Val Tyr Tyr Cys Met Gln His Ile Glu Tyr Pro Phe Thr Phe Gly
158 245 250 255
161 Gln Gly Thr Lys Leu Glu Ile Lys Gly Gly Gly Gly Ser Gly Gly Gly
162 260 265 270
165 Gly Ser Gly Gly Gly Gly Ser Gln Val Gln Leu Val Gln Ser Gly Pro
166 275 280 285

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169 Glu Val Lys Lys Pro Gly Ala Ser Val Lys Val Ser Cys Lys Ala Ser
170      290      295      300
173 Gly Tyr Thr Phe Thr Asn Ser Trp Met Asn Trp Val Arg Gln Arg Pro
174 305      310      315      320
177 Gly Lys Gly Leu Glu Trp Met Gly Arg Ile Tyr Pro Gly Asp Gly Glu
178      325      330      335
181 Thr Ile Tyr Asn Gly Lys Phe Arg Val Arg Val Thr Ile Thr Ala Asp
182      340      345      350
185 Glu Ser Thr Ser Thr Ala Tyr Met Glu Leu Ser Ser Leu Arg Ser Glu
186      355      360      365
190 Asp Thr Ala Val Tyr Tyr Cys Ala Arg Gly Tyr Asp Asp Tyr Ser Phe
191      370      375      380
194 Ala Tyr Trp Gly Gln Gly Thr Thr Val Thr Val Ser Ser Gly Gly Gly
195 385      390      395      400
198 Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Asp Ile Val Met
199      405      410      415
202 Thr Gln Ser Ala Leu Ser Leu Pro Val Thr Pro Gly Glu Pro Ala Ser
203      420      425      430
206 Ile Ser Cys Arg Ser Ser Lys Ser Leu Leu His Ser Asn Gly Asn Thr
207      435      440      445
210 Tyr Leu Tyr Trp Phe Gln Gln Lys Pro Gly Gln Ser Pro Gln Leu Leu
211      450      455      460
214 Ile Tyr Arg Met Ser Asn Leu Ala Ser Gly Val Pro Asp Arg Phe Ser
215 465      470      475      480
218 Gly Ser Gly Ser Gly Thr Ala Phe Thr Leu Lys Ile Ser Arg Val Glu
219      485      490      495
222 Ala Glu Asp Val Gly Val Tyr Tyr Cys Met Gln His Ile Glu Tyr Pro
223      500      505      510
226 Phe Thr Phe Gly Gln Gly Thr Lys Leu Glu Ile Lys
227      515      520
230 <210> SEQ ID NO: 3
231 <211> LENGTH: 5
232 <212> TYPE: PRT
233 <213> ORGANISM: Mus musculus
235 <400> SEQUENCE: 3
237 Ser Ser Trp Met Asn
238 1      5
241 <210> SEQ ID NO: 4
242 <211> LENGTH: 17
243 <212> TYPE: PRT
244 <213> ORGANISM: Mus musculus
246 <400> SEQUENCE: 4
248 Arg Thr Tyr Pro Gly Asp Gly Asp Thr Asn Tyr Asn Gly Lys Phe Lys
249 1      5      10      15
252 Gly
256 <210> SEQ ID NO: 5
257 <211> LENGTH: 13
258 <212> TYPE: PRT
259 <213> ORGANISM: Mus musculus

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261 <400> SEQUENCE: 5
263 Gly Trp Ile Leu Ala Asp Gly Gly Tyr Ser Phe Ala Tyr
264 1 5 10
267 <210> SEQ ID NO: 6
268 <211> LENGTH: 5
269 <212> TYPE: PRT
270 <213> ORGANISM: Mus musculus
272 <400> SEQUENCE: 6
274 Ser Ser Trp Met Asn
275 1 5
278 <210> SEQ ID NO: 7
279 <211> LENGTH: 17
280 <212> TYPE: PRT
281 <213> ORGANISM: Mus musculus
283 <400> SEQUENCE: 7
285 Arg Ile Tyr Pro Gly Asp Gly Asp Thr Asn Tyr Asn Gly Lys Phe Lys
286 1 5 10 15
289 Gly
293 <210> SEQ ID NO: 8
294 <211> LENGTH: 9
295 <212> TYPE: PRT
296 <213> ORGANISM: Mus musculus
298 <400> SEQUENCE: 8
300 Gly Tyr Ala Asp Tyr Ser Phe Ala Tyr
301 1 5
304 <210> SEQ ID NO: 9
305 <211> LENGTH: 5
306 <212> TYPE: PRT
307 <213> ORGANISM: Mus musculus
309 <400> SEQUENCE: 9
311 Ser Ser Trp Met Asn
312 1 5
315 <210> SEQ ID NO: 10
316 <211> LENGTH: 17
317 <212> TYPE: PRT
318 <213> ORGANISM: Mus musculus
320 <400> SEQUENCE: 10
322 Arg Ile Tyr Pro Gly Asp Gly Glu Thr Asn Tyr Asn Gly Lys Phe Lys
323 1 5 10 15
326 Gly
330 <210> SEQ ID NO: 11
331 <211> LENGTH: 9
332 <212> TYPE: PRT
333 <213> ORGANISM: Mus musculus
335 <400> SEQUENCE: 11
337 Gly Phe Gly Asp Tyr Ser Phe Ala Tyr
338 1 5
341 <210> SEQ ID NO: 12
342 <211> LENGTH: 5

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Input Set : A:\14875-153US1sq.txt

Output Set : N:\CRF4\10142005\J551504.raw

343 <212> TYPE: PRT
344 <213> ORGANISM: Mus musculus
346 <400> SEQUENCE: 12
348 Ser Ser Trp Met Asn
349 1 5
352 <210> SEQ ID NO: 13
353 <211> LENGTH: 17
354 <212> TYPE: PRT
355 <213> ORGANISM: Mus musculus
357 <400> SEQUENCE: 13
359 Arg Ile Tyr Pro Gly Asp Gly Asp Thr Asn Tyr Asn Gly Lys Phe Lys
360 1 5 10 15
363 Gly
367 <210> SEQ ID NO: 14
368 <211> LENGTH: 9
369 <212> TYPE: PRT
370 <213> ORGANISM: Mus musculus
372 <400> SEQUENCE: 14
374 Gly Tyr Ala Asp Tyr Ser Phe Ala Tyr
375 1 5
378 <210> SEQ ID NO: 15
379 <211> LENGTH: 5
380 <212> TYPE: PRT
381 <213> ORGANISM: Mus musculus
383 <400> SEQUENCE: 15
385 Arg Ser Trp Met Asn
386 1 5
389 <210> SEQ ID NO: 16
390 <211> LENGTH: 17
391 <212> TYPE: PRT
392 <213> ORGANISM: Mus musculus
394 <400> SEQUENCE: 16
396 Arg Ile Tyr Pro Gly Asp Gly Asp Thr Asn Tyr Asn Gly Lys Phe Lys
397 1 5 10 15
400 Gly
404 <210> SEQ ID NO: 17
405 <211> LENGTH: 9
406 <212> TYPE: PRT
407 <213> ORGANISM: Mus musculus
409 <400> SEQUENCE: 17
411 Gly Tyr Asp Asp Tyr Ser Phe Ala Tyr
412 1 5
415 <210> SEQ ID NO: 18
416 <211> LENGTH: 5
417 <212> TYPE: PRT
418 <213> ORGANISM: Mus musculus
420 <400> SEQUENCE: 18
422 Asn Ser Trp Met Asn
423 1 5

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/551,504

DATE: 10/14/2005
TIME: 10:55:56

Input Set : A:\14875-153US1sq.txt
Output Set: N:\CRF4\10142005\J551504.raw

Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete,
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:166,167,168,169,170,171,172,173,174,175,176,177,178,179,180,181,182,183

Seq#:184,185,186,187,188,189,190,191,192,193,194,195,196,197,198,199,200,201

Seq#:202,203,204,205,206,207,208,209,210,211,212,213,214,215,216,217,218,219

Seq#:220,221,222,223,224,225,226,227

VERIFICATION SUMMARY

DATE: 10/14/2005

PATENT APPLICATION: US/10/551,504

TIME: 10:55:56

Input Set : A:\14875-153US1sq.txt

Output Set: N:\CRF4\10142005\J551504.raw

L:13 M:270 C: Current Application Number differs, Replaced Current Application No

L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date